

SEQUENCE LISTING

<110> Yuko, Wakamatsu and Kenjiro, Ozato

<120> See-Through Medaka

<130> PH-1134

<140>

<141>

<150> JP 2000/172375

<151> 2000-06-08

<160> 8

<170> PatentIn Ver. 2.0

<210> 1

<211> 25

<212> DNA

<213> *Oryzias latipes*

<220>

<223> Other information: Forward primer pHO5.5-F.

<400> 1

cctgcaatgg gaaattattc tgctc

25

<210> 2

<211> 28

<212> DNA

<213> *Oryzias latipes*

<220>

<223> Other information: Reverse primer pHO5.5-RV.

<400> 2

cttttggtgc ttggttatg aaacgatg

28

<210> 3

<211> 19

<212> DNA

<213> Unknown

<220>

<223> Other information: Common sequence found in animals. The sequence is used herein as a forward primer to amplify the short fragment of vasa cDNA.

<400> 3

atggcntgyg encaracng

19

<210> 4

<211> 18

<212> DNA

<213> Unknown

<220>

<223> Other information: Common sequence found in animals. The sequence is used herein as a reverse primer used to amplify the short fragment of vasa cDNA.

<400> 4

raancccatr tcyaacat

18

<210> 5

<211> 20

<212> DNA

<213> Oryzias latipes

<220>

<223> Other information: Primer T7 used to amplify the 3' region of vasa gene.

<400> 5

taatacgact cactataggg

20

<210> 6

<211> 23

<212> DNA

<213> *Oryzias latipes*

<220>

<223> Other information: Primer VI-8 used to amplify the 3' region of vasa gene.

<400> 6

aggaggtgcc gtcattgctg gag

23

<210> 7

<211> 30

<212> DNA

<213> *Oryzias latipes*

<220>

<223> Other information: Primer VP1M used to amplify 5.1 kb genomic fragment comprising the promotor region of vasa gene.

<400> 7

cctcccagtc gtccatatga atcgtctgat

30

<210> 8

<211> 37

<212> DNA

<213> *Oryzias latipes*

<220>

<223> Other information: Primer VP3 used to amplify 5.1 kb genomic fragment comprising the promotor region of vasa gene.

<400> 8

agaggatcca atagaatgag taatggttct ctatttc

37